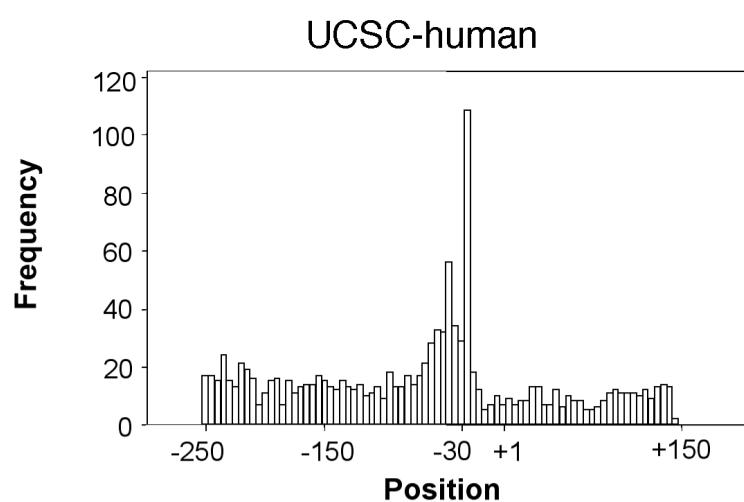


A



B

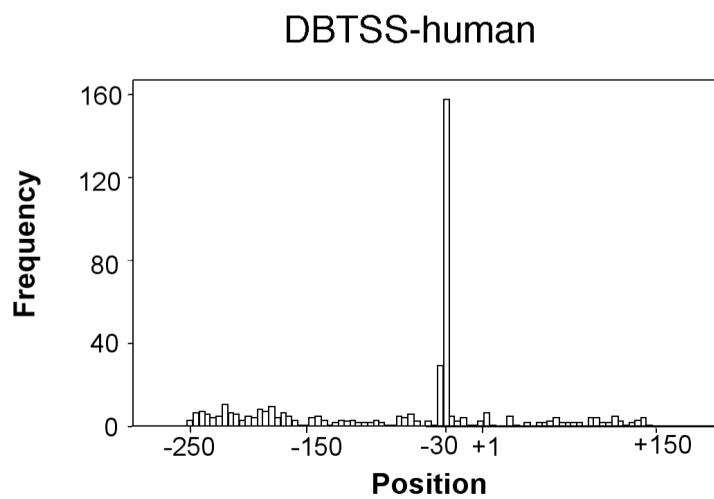


Fig. S3. Comparison of distribution of TATA-8 sequences in the human genes in the UCSC and DBTSS databases.

The 15,685 non redundant, annotated human genes in the UCSC database and the 10,271 non redundant, annotated human genes in the DBTSS database were searched for the 8-best TATA sequences (see Fig. 1 main text) in the -250 to +150 bp region relative to the transcription start site (+1).